OIPE

#3

RAW SEQUENCE LISTING   DATE: 02/10/2002
PATENT APPLICATION: US/09/954,773A   TIME: 13:24:59

Input Set : **A:\EP.txt**
Output Set: N:\CRF3\02102002\I954773A.raw

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   3  <110> APPLICANT: Southern Illinois University
   4        Lightfoot, David A
   5        Gibson, Paul T
   6        Meksem, Khalid
   8  <120> TITLE OF INVENTION: SOYBEAN SUDDEN DEATH SYNDROME RESISTANT SOYBEANS, SOYBEAN
CYST NEMATODE
   9        RESISTANT SOYBEANS AND METHODS OF BREEDING AND IDENTIFYING RESISTANT PLANTS
  11  <130> FILE REFERENCE: 1268/2/2
C--> 13 <140> CURRENT APPLICATION NUMBER: US/09/954,773A
C--> 13 <141> CURRENT FILING DATE: 2002-01-22
  13  <150> PRIOR APPLICATION NUMBER: 09/007,119
  14  <151> PRIOR FILING DATE: 1998-01-14
  16  <160> NUMBER OF SEQ ID NOS: 20
  18  <170> SOFTWARE: PatentIn version 3.1
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  22  <212> TYPE: DNA
  23  <213> ORGANISM: Glycine max
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  27  <222> LOCATION: (1)..(527)
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  36  ttttattaaa taataaaaga gtaattacat aaaaatatgt tcgattacat taaaccccaa     180
  38  caaaggatga atttagcttc tcatgaccat ggggaaaatc aaacttgatg aacaagaaga     240
  40  tgaagaagaa tccttaagga taaacactgc ctagctccaa tgtgctctct agtattttat     300
  42  ctttcaaaaa tccccaagaa cccctaattt tcagtaagaa gcccattttc aatcagaagc     360
  44  ccattttcaa tcaagaagcc cattttcaat cagaagccca ttttcaatca gaagcccatt     420
  46  ttcaatcaga gcccatttt caatcagaag cccattttat aattgtattc ccaaaacttg     480
  48  agattcttga acgtaaatta ttagtaaatt gtaatcacct ctgtaaa                  527
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  59  <223> OTHER INFORMATION: n is a, c, g, or t/u
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  67  aaggtaacag caataagtaa tcccttgttt ataagatccc agaacttcca gtttatttaa     180
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         73 ttacctatgt tgttccacaa gaatatactc aaatgacttt gccttgattt aaattatcac     360
W--> 75 gatgtaacac aaacaaagat gatantttgt cgatcaactg ttcagcacca agagagccct     420
W--> 77 ccccacaatc aactcaggtt ttcacttttg gtgcttgaaa atgagtggca catgnaaaag     480
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W--> 81 tgagccaaaa catcatctgg gncatcaacc ttcatgnctt tntcaagttt atacctatna     600
W--> 83 ntnactangt cttatatttn canntggtga ttacanttac nantaagttt agcttnaaga     660
W--> 85 aatncaagtt ttngggactc catgcctngn cnggntttcn natccgtcgg ccagggcggn     720
W--> 87 cnggnncact gntnggnagn cccanttncn cagancacng ncccntttcc attccnggnc     780
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        108 gtttatagta acttcattgc aaaactcaac aaatagattt ttggaacctt aatataataa     180
        110 aattcaacag tcttctttaa ttttattctg ctcttacctt ctcataggat catatagaat     240
        112 ttaaccctac aagctctcaa aaaacaatcc attattatgc tccttatcca ataaaacaaa     300
W--> 114 accatagagt gattctcaaa atgaagattg acaaaggcaa aaagttatgc tggntcaata     360
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        129 <223> OTHER INFORMATION: n is a, c, g, or t/u
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        139 agt                                                                   183
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        158 tgttttatga aggtaataca ccaattatta tggttttttg tttaataaaa tgtgaataat   180
        160 tgtcaatcgt gattgcatta tctctccttt actctgtctc ttcacctttt ttacccttt    240
        162 atttgagagg aagaatccat gtagtaaaaa atgatgataa aattgttaga aaatatagtg   300
        164 tcatgtaatt agagattcag attataactt agaagacact attattttca tgtaatacta   360
        166 tccacgggta attatcaata ctgacatatt ttcactcaaa atattctggt tttctcatta   420
W--> 168 tatacattta aataggagct attanccatt gcaagcttgg gtttggaggc cttccgatgc   480
W--> 170 cttgttggga ttgngacca                                                  499
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        174 <211> LENGTH: 500
        175 <212> TYPE: DNA
        176 <213> ORGANISM: Glycine max
        178 <220> FEATURE:
        179 <221> NAME/KEY: misc_feature
        180 <222> LOCATION: (1)..(500)
        181 <223> OTHER INFORMATION: n is a, c, g, or t/u
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W--> 187 taaattgaaa atatatacgt aagancttca tctaacagtg ctagtcgaag aatgcgtaaa   120
W--> 189 tgcaggnnat ccatttccat actaaaatgg acaaaactta tatttttttt ttagcggcaa   180
W--> 191 acgttaatta ttaatttttt ttagtacaag ggatcaaacc angacctttc ccttctttcc   240
        193 atctttcttg accacccaac caaccttata tctccacaaa acttattata tgttgttctt   300
W--> 195 cggggactat cagaattgga gtttaacctc gggcantcaa tctacataat ccttgatttn   360
W--> 197 atttngtgaa gttctaaagc cacaggcatt atttatntta ttntttctgn agtaacccnc   420
W--> 199 catatgttgg tnnataaggg tangnatnaa aatncnttgg ntggtnncna tttgcncttn   480
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        207 <213> ORGANISM: Glycine max
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        210 <221> NAME/KEY: misc_feature
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        212 <223> OTHER INFORMATION: n is a, c, g, or t/u
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W--> 220 gttcattgta gcaatagcat cgttatctct tattattgct ggttcatcat natctcaatt   180
        222 ccagtggca                                                            189
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        228 <213> ORGANISM: Glycine max
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        232 <222> LOCATION: (1)..(724)
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RAW SEQUENCE LISTING   DATE: 02/10/2002
PATENT APPLICATION: US/09/954,773A   TIME: 13:24:59

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241 taacttatta acgtttcaca ttttcaaaga cgagtctagc tatttataat ttttttttcct    180
243 aaaatatatt ttttgtcctc ataaatatga aaatatttaa aattcgttcc taattttttt    240
245 ttcaaagcat cttccttct cacaaaattg aaatgtatca ttttttttttg ttcaaaagtt    300
247 taaataaatt tgaacctaat atgacatttt atatcggtta tacatataac tgatataaac    360
249 atcaagtttt ttatatcaat gatacctata actgatatca aatgtgacaa ttatatatat    420
251 aattaatgta aaaaagtcat aaatataatt tattttgagt caaaaaataa tatattttaa    480
W--> 253 ttattttgaa gatgaaaaag gataaattta aaacatttgt gtgangatga aaaactagat    540
255 gtttttttttc ctggtttaaa tgcaaaacca atgctatttt atttaaattt tacctttttt    600
W--> 257 ttataattac nccaccaaaa aaccgtttgg tgttacaaat ttganttaaa ttctnttgtt    660
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261 atct                                                                724
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265 <211> LENGTH: 801
266 <212> TYPE: DNA
267 <213> ORGANISM: Glycine max
269 <220> FEATURE:
270 <221> NAME/KEY: misc_feature
271 <222> LOCATION: (1)..(801)
272 <223> OTHER INFORMATION: n is a, c, g, or t
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280 gcacacaccc tatctttcat aaaattacta cacttttttaa tttttgtaat aaaaaaccta    180
282 gaaaaactca ttatgaaaca gatgatgtac ttttaacact ctgtcggcct ctctctctct    240
284 attatatatt gatttaaatt tattgagaat tatattttttg ttgggtctca tttattatat    300
286 tttattaatt ggatccgggc cctctagatg cggccgcatg cataagcttg agtattctat    360
288 agtgtcacct aaatagcttg gcgtaatcat ggtcatagct gtttcctgtg tgaaattgtt    420
W--> 290 atccgctcac aattccacac aacatacgag ccggaagcat aaagtgtnaa gcctggggtn    480
W--> 292 cctaatgagt gagctaactc acattaattg ccttgcgctc actgcccgct ttccagtcng    540
W--> 294 gaaacctgtc ctgccagctg cattaatgaa tcngccaacc cncggggana agcngtttgc    600
W--> 296 ntatgggcgc tcttnccgct tcctcgctca ntgactcgct gcgctcngtc nttcngntgc    660
W--> 298 cgcgaacggt atcancncac tcnaangnng taaatacggt tatccaccna accnngggga    720
W--> 300 naacccngga aaaacatgt nanccaaaag gccnccaaaa ggccangaaa cnttnaaaag    780
W--> 302 gcccnnttgc ttgncttttnt n                                             801
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307 <212> TYPE: DNA
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310 <220> FEATURE:
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313 <223> OTHER INFORMATION: n is a, c, g, or t
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321 tcttgtactt ttaacttttta agtcatactc cctttggact catatataag caaaagagtg    180
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        327 atgataacaa gttccaatga aggacatttt agaaataacc ttatttttta tttgagatta    360
        329 gtaaaattaa atgatgtgaa ctaactttct taattaatgt gaaattagtt attttttctt    420
        331 atatacgagt ccaaagggag taccaaattt cacaaatgta ctaaatgta ttatatgctt     480
W--> 333 cttttaatt catctttgct gcatanctac ttagctactg tgctctgatc cgggccctct        540
        335 agatgcggcc gcatgcataa gcttgagtat ctatagtgtc cctaaatagc ttggcgtatc    600
W--> 337 atggtcatag ctgtttccng tgtgaaattg ttatccgctc acaattccac acaacatacg       660
W--> 339 anccggaagc ataaaagtgt taagccnggg gtgcctaatg agtgagctaa ctcacattaa       720
W--> 341 ttgcgttgcg ctcactgccc gcttccnatt cgggaaactg tcctgncanc tgcattaatg       780
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        362 agagggctg attttggaga aaacatcatc catggtataa agtccgttta gattccagct      180
        364 attgttcaca ttcatccctt acatatgaga atatccctat aagctgaaac taactttttac  240
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        368 ttcagaaatt cacataaata actaaaagta agttgtattt tatttatgtc tggatttact     360
        370 gcacaaacta aacaaaagtt tgtggattta gacataaaaa ataccaatgc tgtgtgaaaa     420
W--> 372 taagaaatgg tggtcatata gacaagtttc ttttctgttt tctttaaatt gcagtcnaag      480
W--> 374 ccatcangag gttcatgtaa ttaaccaaac tagacgttga cttttggttt tatccttttg      540
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W--> 378 cgcaaatttg ttaaaatatt naataataat atcacctaaa atcatatttg tcanttcatt      660
W--> 380 ttgtttttang ttatatcaat tattatttt tacctttacnt cctttataat ntcaatgatg     720
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        403 caaattcttt aatgaaaagt taattacata aaatatttta gtagaagcaa ttttacacag    180
        405 ttattatta aaaaaattac acagttattc aataacaaat tacaatatat tataaggtta      240
        407 taataaatat tttaaaattc atataaaaga tgacttatta ataagttgat aatgtaaatt    300
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VERIFICATION SUMMARY   DATE: 02/10/2002
PATENT APPLICATION: . US/09/954,773A   TIME: 13:25:00

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L:13 M:271 C: Current Filing Date differs, Replaced Current Filing Date
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PATENT APPLICATION: US/09/954,773A   TIME: 13:25:00

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